Gambel PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/08/477,989B

#191644

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This Raw Listing contains the General Information Section and up to the first 5 pages.

1	SEQUENCE LISTING (1) General Information: (i) APPLICANT: Bazin, Herv Latinne, Dominique Kaplan, Ruth
2	
3	(1) General Information:
4	(i) APPLICANT: Bazin, Herv
[.] 5	Latinne, Dominique
6	Kaplan, Ruth
7	Kieber-Emmons, Thomas
8	Postema, Christina E.
9	White-Scharf, Mary
10	(ii) TITLE OF INVENTION: LO-CD2a Antibody and Uses
11	Thereof for Inhibiting
12	T-Cell Activation and
13	Proliferation
14	(iii) NUMBER OF SEQUENCES: 96
15	(iv) CORRESPONDENCE ADDRESS:
16	(A) ADDRESSEE: Carella, Byrne, Bain, Gilfillan,
17	Cecchi, Stewart & Olstein
18	(B) STREET: 6 Becker Farm Road
19	(C) CITY: Roseland
20	(D) STATE: New Jersey
21	(E) COUNTRY: U.S.A.
22	(F) ZIP: 07068
23	
24	(V) COMPUTER READABLE FORM:
25	(A) MEDIUM TYPE: 3.5 inch diskette
26	(B) COMPUTER: IBM PS/2
27	(C) OPERATING SYSTEM: MS-DOS
28	(D) SOFTWARE: WordPerfect 5.1
29	
30	(vi) CURRENT APPLICATION DATA:
31	(A) APPLICATION NUMBER: 08/477,989
32	(B) FILING DATE: 07-JUN-1995
33	(C) CLASSIFICATION:
34	
-35	(Vii) PRIOR APPLICATION DATA:
36	(A) APPLICATION NUMBER: 08/407,009
37	(B) FILING DATE: 29-MAR-1995
38	(A) APPLICATION NUMBER: 08/119,032
39	(B) FILING DATE: 09-SEP-1993
40	(A) APPLICATION NUMBER: 08/027,008
41	(B) FILING DATE: 05-MAR-1993
42	
43	(viii) ATTORNEY/AGENT INFORMATION:
44	(A) NAME: Olstein, Elliot M.
45	(B) REGISTRATION NUMBER: 24,025
46	(C) REFERENCE/DOCKET NUMBER: 61750-147

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47
               (ix) TELECOMMUNICATION INFORMATION:
48
                     (A) TELEPHONE: 201-994-1700
49
                     (B) TELEFAX: 201-994-1744
50
51
52
    (2) INFORMATION FOR SEQ ID NO: 1:
53
               (i) SEQUENCE CHARACTERISTICS:
                     (A) LENGTH: 32 bases
54
                     (B) TYPE: nucleic acid
55
                     (C) STRANDEDNESS: single
56
                     (D) TOPOLOGY: linear
57
               (ii) MOLECULE TYPE: oligonucleotide
58
59
               (ix) FEATURE:
60
61
                     (A) NAME/KEY: PCR primer
62
               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                      32
    AACCCGGGGA CATTCAGCTG ACCCAGTCTC AA
63
64
65
               (2) INFORMATION FOR SEQ ID NO: 2:
               (i) SEQUENCE CHARACTERISTICS:
66
                     (A) LENGTH: 29 bases
67
                     (B) TYPE: nucleic acid
68
69
                     (C) STRANDEDNESS: single
70
                     (D) TOPOLOGY: linear
71
               (ii) MOLECULE TYPE: oligonucleotide
72
               (ix) FEATURE:
73
74
                     (A) NAME/KEY: PCR primer
               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
75
    CAGTCGACTA CAGTTGGTGC AGCATCAGC
                                                                      29
76
77
               (2) INFORMATION FOR SEQ ID NO: 3:
78
79
               (i) SEQUENCE CHARACTERISTICS:
80
                     (A) LENGTH: 31 bases
81
                     (B) TYPE: nucleic acid
                     (C) STRANDEDNESS: single
82
               (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide
83
84
85
86
               (ix) FEATURE:
87
                     (A) NAME/KEY: PCR primer
               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
88
    AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G
89
                                                                      31
90
91
               (2) INFORMATION FOR SEQ ID NO: 4:
92
               (i) SEQUENCE CHARACTERISTICS:
93
                     (A) LENGTH: 27 bases
94
                     (B) TYPE: nucleic acid
95
                     (C) STRANDEDNESS: single
96
                     (D) TOPOLOGY: linear
               (ii) MOLECULE TYPE: oligonucleotide
97
98
99
               (ix) FEATURE:
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100	(A) NAME/KEY: PCR primer	
101	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
.102	AAGTCGACCC AGTGGATAGA CCGATGG	27
103	(A) THEODYNATON BOD GEO TO NO. F.	
104	(2) INFORMATION FOR SEQ ID NO: 5:	
105	(i) SEQUENCE CHARACTERISTICS:	
106	(A) LENGTH: 22 bases (B) TYPE: nucleic acid	
107	(C) STRANDEDNESS: single	
108 109	(D) TOPOLOGY: linear	
110	(ii) MOLECULE TYPE: oligonucleotide	
111	(11) Moddeodd 111d. Ollgondoleocide	
112	(ix) FEATURE:	
113	(A) NAME/KEY: PCR primer	
114	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
115	CCGCAAGCTT CATGGGATGG AG	22
116		
117	(2) INFORMATION FOR SEQ ID NO: 6:	
118	(i) SEQUENCE CHARACTERISTICS:	
119	(A) LENGTH: 27 bases	
120	(B) TYPE: nucleic acid	
121	(C) STRANDEDNESS: single	
122	(D) TOPOLOGY: linear	
123	(ii) MOLECULE TYPE: oligonucleotide	
124		
125	(ix) FEATURE:	
126	(A) NAME/KEY: PCR primer	
127	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
128 129	GCTGCTTGGG GACTGGGTCA GCTGGAT	27
130	(2) INFORMATION FOR SEQ ID NO: 7:	
131	(i) SEQUENCE CHARACTERISTICS:	
132	(A) LENGTH: 21 bases	
133	(B) TYPE: nucleic acid	
134	(C) STRANDEDNESS: single	
135	(D) TOPOLOGY: linear	
136	(ii) MOLECULE TYPE: oligonucleotide	
137	•	
138	(ix) FEATURE:	
139	(A) NAME/KEY: PCR primer	
140	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
141	ATTCAGCTGA CCCAGTCTCC A	21
142		
143	(2) INFORMATION FOR SEQ ID NO: 8:	
144	(i) SEQUENCE CHARACTERISTICS:	
145	(A) LENGTH: 58 bases	
146	(B) TYPE: nucleic acid	
147	(C) STRANDEDNESS: single	
148	(D) TOPOLOGY: linear	
149 150	(ii) MOLECULE TYPE: oligonucleotide	
150	(ix) FEATURE:	
152	(A) NAME/KEY: PCR primer	
132	(A) NAME/REI: FCR PLIMET	

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	INPUI SEI: SZ/
153	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
154	GATCGGATCC ACCTGAGGAA GCAAAGTTTA AATTCTACTC ACGTTTCAGT TCCAGCTT58
155	(2) INFORMATION FOR SEQ ID NO: 9:
156 157	(i) SEQUENCE CHARACTERISTICS:
157	(A) LENGTH: 31 bases
159	(B) TYPE: nucleic acid
160	(C) STRANDEDNESS: single
161	(D) TOPOLOGY: linear
162	(ii) MOLECULE TYPE: oligonucleotide
163	(11) Hollood III at Oligonad Cooled
164	(ix) FEATURE:
165	(A) NAME/KEY: PCR primer
166	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
167	TCTCCTGCAG TGGGACCTCG GAGTGGACAC C 31
168	
169	(2) INFORMATION FOR SEQ ID NO: 10:
170	(i) SEQUENCE CHARACTERISTICS:
171	(A) LENGTH: 21 bases
172	(B) TYPE: nucleic acid
173	(C) STRANDEDNESS: single
174	(D) TOPOLOGY: linear
175	(ii) MOLECULE TYPE: oligonucleotide
176	
177	(ix) FEATURE:
178	(A) NAME/KEY: PCR primer
179	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
180	GAGGTCCAGC TGCAGCAGTC T 21
181	(A) THROPHIMTON FOR GEO TO NO. 11.
182 183	(2) INFORMATION FOR SEQ ID NO: 11:
184	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 bases
185	(B) TYPE: nucleic acid
186	(C) STRANDEDNESS: single
187	(D) TOPOLOGY: linear
188	(ii) MOLECULE TYPE: oligonucleotide
189	(, <u></u>
190	(ix) FEATURE:
191	(A) NAME/KEY: PCR primer
192	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
193	CGATGTATCA GCTGTCAGTG TGGC 24
194	
195	(2) INFORMATION FOR SEQ ID NO: 12:
196	(i) SEQUENCE CHARACTERISTICS:
197	(A) LENGTH: 24 bases
198	(B) TYPE: nucleic acid
199	(C) STRANDEDNESS: single
200	(D) TOPOLOGY: linear
201	(ii) MOLECULE TYPE: oligonucleotide
202	(Am) BRADITOR
203	(ix) FEATURE:
204 205	(A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
405	(XI) SEGOUNCE DESCRIPTION: SEG ID NO: 15:

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		1111 01 021. 02/2/
206	GCCACACTGA CAGCTGATAC ATCG	24
207		
208	(2) INFORMATION FOR SEQ ID NO: 13:	
209	(i) SEQUENCE CHARACTERISTICS:	
210	(A) LENGTH: 21 bases	
211	(B) TYPE: nucleic acid	
212	(C) STRANDEDNESS: single	
213	(D) TOPOLOGY: linear	
214	(ii) MOLECULE TYPE: oligonucleotide	
215		
216	(ix) FEATURE:	
217	(A) NAME/KEY: PCR primer	
218	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:	
219	CAGAGTGCCT TGGCCCCAGT A	21
220		
221	(2) INFORMATION FOR SEQ ID NO: 14:	
222	(i) SEQUENCE CHARACTERISTICS:	
223	(A) LENGTH: 27 bases	
224	(B) TYPE: nucleic acid	
225	(C) STRANDEDNESS: single	
226	(D) TOPOLOGY: linear	
227	(ii) MOLECULE TYPE: oligonucleotide	
228		
229	(ix) FEATURE:	
230	(A) NAME/KEY: PCR primer	
231	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:	•
232	TACTGGGGCC AAGGCACCCT CGTCACA	27
233	,	
234	(2) INFORMATION FOR SEQ ID NO: 15:	
235	(i) SEQUENCE CHARACTERISTICS:	
236	(A) LENGTH: 25 bases	
237	(B) TYPE: nucleic acid	
238	(C) STRANDEDNESS: single	
239	(D) TOPOLOGY: linear	
240	(ii) MOLECULE TYPE: oligonucleotide	
241	(, __	
242	(ix) FEATURE:	
243	(A) NAME/KEY: PCR primer	
244	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:	
245	GATCGGATCC CTATAAATCT CTGGC	25
246		
247	(2) INFORMATION FOR SEQ ID NO: 16:	
248	(i) SEQUENCE CHARACTERISTICS:	
249	(A) LENGTH: 43 bases	
250	(B) TYPE: nucleic acid	
251	(C) STRANDEDNESS: single	
252	(D) TOPOLOGY: linear	
253	(ii) MOLECULE TYPE: oligonucleotide	
254	(11) Hondoon III D. Offgondorootide	
255	(ix) FEATURE:	
256	(A) NAME/KEY: PCR primer	
257	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:	
258	TTGGATCCGC GGCCGCGTCG ACTACAGTTG GTGCAGCATC AGC	43
	110001000 000000100 AULAOAUITO UIOCAUCATO AUC	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/477,989B*

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